

RAW SEQUENCE LISTING

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Application Serial Number: 09/993,292C
Source: IFW/6
Date Processed by STIC: 7/15/05

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IFW16

RAW SEQUENCE LISTING

DATE: 07/15/2005

PATENT APPLICATION: US/09/993,292C

TIME: 10:04:59

Input Set : A:\09-993,292 Sequence Listing.txt

Output Set: N:\CRF4\07152005\I993292C.raw

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3 <110> APPLICANT: University of Maryland, Baltimore
4   GALEN, James E.
6 <120> TITLE OF INVENTION: USE OF CLYA HEMOLYSIN FOR EXCRETION OF PROTEINS
8 <130> FILE REFERENCE: A8461
10 <140> CURRENT APPLICATION NUMBER: 09/993,292C
11 <141> CURRENT FILING DATE: 2001-11-23
13 <150> PRIOR APPLICATION NUMBER: US 60/252,516
14 <151> PRIOR FILING DATE: 2000-11-22
16 <160> NUMBER OF SEQ ID NOS: 25
18 <170> SOFTWARE: PatentIn version 3.3
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 6271
22 <212> TYPE: DNA
23 <213> ORGANISM: Artificial Sequence
25 <220> FEATURE:
26 <223> OTHER INFORMATION: pSEC84 Expression Plasmid
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35 tcagtgtctg caaataactta agaataagtt attgatttta acctgaatt attattgctt    240
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39 ttttgaaaca tctatagcga taaatgaaac atcttaaaaag ttttagtata atattcgtgt    360
41 tggattattc tgcatttttg gggagaatgg acttgccgac tgattaatga gggttaatca    420
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59 aagcgcaaaa atctctcctg acaagttcac aaagtttcaa caacgcttcc ggaaaactgc    960
61 tggcattaga tagccagtta actaatgatt ttccggaaaa aagtagttat ttccagtcac   1020
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89 gtccaacccg gtaagacacg acttatcgcc actggcagca gccactggta acaggattag 1860
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179 gaaaataatc cgctcattca gaccggttcac gggaaatccg tgtgattggt gccgcatcac 4560
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187 aaaaaaacac cataaggagt ttataaatg ttggtattca ttgatgacgg ttcaacaaac 4800
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237 gctctgtcat tttctgaaac tcttcatgct g 6271
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241 <211> LENGTH: 305
242 <212> TYPE: PRT
243 <213> ORGANISM: Salmonella typhi
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248 1 5 10 15
251 Ile Glu Thr Ala Asp Gly Ala Leu Asp Leu Tyr Asn Lys Tyr Leu Asp
252 20 25 30
255 Gln Val Ile Pro Trp Lys Thr Phe Asp Glu Thr Ile Lys Glu Leu Ser
256 35 40 45
259 Arg Phe Lys Gln Glu Tyr Ser Gln Glu Ala Ser Val Leu Val Gly Asp
260 50 55 60
263 Ile Lys Val Leu Leu Met Asp Ser Gln Asp Lys Tyr Phe Glu Ala Thr
264 65 70 75 80
267 Gln Thr Val Tyr Glu Trp Cys Gly Val Val Thr Gln Leu Leu Ser Ala
268 85 90 95
271 Tyr Ile Leu Leu Phe Asp Glu Tyr Asn Glu Lys Lys Ala Ser Ala Gln
272 100 105 110

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275 Lys Asp Ile Leu Ile Arg Ile Leu Asp Asp Gly Val Lys Lys Leu Asn
276      115      120      125
279 Glu Ala Gln Lys Ser Leu Leu Thr Ser Ser Gln Ser Phe Asn Asn Ala
280      130      135      140
283 Ser Gly Lys Leu Leu Ala Leu Asp Ser Gln Leu Thr Asn Asp Phe Ser
284 145      150      155      160
287 Glu Lys Ser Ser Tyr Phe Gln Ser Gln Val Asp Arg Ile Arg Lys Glu
288      165      170      175
291 Ala Tyr Ala Gly Ala Ala Ala Gly Ile Val Ala Gly Pro Phe Gly Leu
292      180      185      190
295 Ile Ile Ser Tyr Ser Ile Ala Ala Gly Val Ile Glu Gly Lys Leu Ile
296      195      200      205
299 Pro Glu Leu Asn Asn Arg Leu Lys Thr Val Gln Asn Phe Phe Thr Ser
300      210      215      220
303 Leu Ser Ala Thr Val Lys Gln Ala Asn Lys Asp Ile Asp Ala Ala Lys
304 225      230      235      240
307 Leu Lys Leu Ala Thr Glu Ile Ala Ala Ile Gly Glu Ile Lys Thr Glu
308      245      250      255
311 Thr Glu Thr Thr Arg Phe Tyr Val Asp Tyr Asp Asp Leu Met Leu Ser
312      260      265      270
315 Leu Leu Lys Gly Ala Ala Lys Lys Met Ile Asn Thr Cys Asn Glu Tyr
316      275      280      285
319 Gln Gln Arg His Gly Lys Lys Thr Leu Phe Glu Val Pro Asp Val Ala
320      290      295      300
323 Ser
324 305
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328 <211> LENGTH: 102
329 <212> TYPE: DNA
330 <213> ORGANISM: Artificial Sequence
332 <220> FEATURE:
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338 tagttaaaag cgcgatcgaa accgcagatg gggcattaga tc      102
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342 <211> LENGTH: 101
343 <212> TYPE: DNA
344 <213> ORGANISM: Artificial Sequence
346 <220> FEATURE:
347 <223> OTHER INFORMATION: Cloning Primer
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356 <211> LENGTH: 97
357 <212> TYPE: DNA
358 <213> ORGANISM: Artificial Sequence
360 <220> FEATURE:

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361 <223> OTHER INFORMATION: Cloning Primer
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370 <211> LENGTH: 69
371 <212> TYPE: DNA
372 <213> ORGANISM: Artificial Sequence
374 <220> FEATURE:
375 <223> OTHER INFORMATION: Cloning Primer
377 <400> SEQUENCE: 6
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380 cgcaacgcg      69
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386 <213> ORGANISM: Artificial Sequence
388 <220> FEATURE:
389 <223> OTHER INFORMATION: Cloning Primer
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396 <211> LENGTH: 101
397 <212> TYPE: DNA
398 <213> ORGANISM: Artificial Sequence
400 <220> FEATURE:
401 <223> OTHER INFORMATION: Cloning Primer
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411 <212> TYPE: DNA
412 <213> ORGANISM: Artificial Sequence
414 <220> FEATURE:
415 <223> OTHER INFORMATION: Cloning Primer
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418 cgaaatagac agatcgctga gataggtgcc tcaactgatta agcattggta atgacctagg      60
420 ctactcatg tttgacagct tatcatcgat aacctttaat g      101
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425 <212> TYPE: DNA
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429 <223> OTHER INFORMATION: Cloning Primer
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VERIFICATION SUMMARY

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